

Supplemental Methods

Mouse QPCR primers:

Col1a1

F: CGATGGATTCCCGTTTCGAGT

R: CGATCTCGTTGGATCCCTGG

Col1a2

F: CTGGTCCTGTTGGAAGTCGT

R: CAGATGCACCTGTTTCTCCA

Mmp1

F: AACTACATTTAGGGGAGAGGTGT

R: GCAGCGTCAAGTTTAACTGGAA

Mmp2

F: CAAGTTCCCCGGCGATGTC

R: TTCTGGTCAAGGTCACCTGTC

Mmp3

F: ACATGGAGACTTTGTCCCTTTTG

R: TTGGCTGAGTGGTAGAGTCCC

Mmp7

F: CTGCCACTGTCCCAGGAAG

R: GGGAGAGTTTTCCAGTCATGG

Mmp8

F: TCTTCCTCCACACACAGCTTG

R: CTGCAACCATCGTGGCATTG

Mmp9

F: CTGGACAGCCAGACACTAAAG

R: CTCGCGGCAAGTCTTCAGAG

Mmp13

F: CTTCTTCTTGTTGAGCTGGACTC

R: CTGTGGAGGTCACCTGTAGACT

Mmp14

F: CAGTATGGCTACCTACCTCCAG

R: GCCTTGCCTGTCACCTTGTAAG

Timp1

F: GCAACTCGGACCTGGTCATAA

R: CGGCCCGTGATGAGAACT

Timp2

F: TCAGAGCCAAAGCAGTGAGC

R: GCCGTGTAGATAAACTCGATGTC

Timp3

F: CTTCTGCAACTCCGACATCGT

R: GGGGCATCTTACTGAAGCCTC

Timp4

F: TGTCTACACGCCATTTGACTC

R: TGGACATCTCCTTACTTGGCA

Ddr1

F: ATGCTGACATGAAGGGACATTT

R: GGTGTAGCCTACGAAGGTCCA

Ddr2

F: ATCACAGCCTCAAGTCAGTGG

R: TTCAGGTCATCGGGTTGCAC

Itga1

F: CCTTCCCTCGGATGTGAGTCA

R: AGTTCTCCCCGTATGGTAAGA

Itga2

F: TGTCTGGCGTATAATGTTGGC

R: CTTGTGGGTTTCGTAAGCTGCT

Itgb1

F: ATGCCAAATCTTGCGGAGAAT

R: TTTGCTGCGATTGGTGACATT

Mfge8

F: AGATGCGGGTATCAGGTGTGA

R: GGGGCTCAGAACATCCGTG

Fap

F: GTCACCTGATCGGCAATTTGT

R: CCCCATCTGAAGGTCGTAGAT

Mrc1

F: CCACTCTGGGCCATGAGGCTTC

R: CTGAATGATCGCATGCTCATTC

Mrc2

F: GAGTCACCCCAGTCTGCAAT

R: CACTGCCATCGAAGACTCAA

Ctsk

F: GAAGAAGACTCACCAGAAGCAG

R: TCCAGGTTATGGGCAGAGATT

Ctsb

F: TCCTTGATCCTTCTTTCTTGCC

R: ACAGTGCCACACAGCTTCTTC

Ctsd

F: GCTTCCGGTCTTTGACAACT

R: CACCAAGCATTAGTTCTCCTCC

Ctsl

F: ATCAAACCTTTAGTGCAGAGTGG

R: CTGTATTCCCCGTTGTGTAGC

Ctss

F: CCATTGGGATCTCTGGAAGAAAA

R: TCATGCCCACTTGGTAGGTAT

Gapdh

F: GTTGTCTCCTGCGACTTCA

R: GGTGGTCCAGGGTTTCTTA

Actb

F: CCAACCGTGAAAAGATGACC

R: ACCAGAGGCATACAGGGACA

Human QPCR Primers:

MRC2

F: CACTGCTATTCTTTCCACAT

R: ACATTCTCCATCTCATCCA

MZF1

F: AGTGTAAGCCCTCACCTCC

R: GGGTCCTGTTCACTCCTCAG

ACTB

F: AGAGCTACGAGCTGCCTGAC

R: AGCACTGTGTTGGCGTACAG

Supplemental Figure Legends

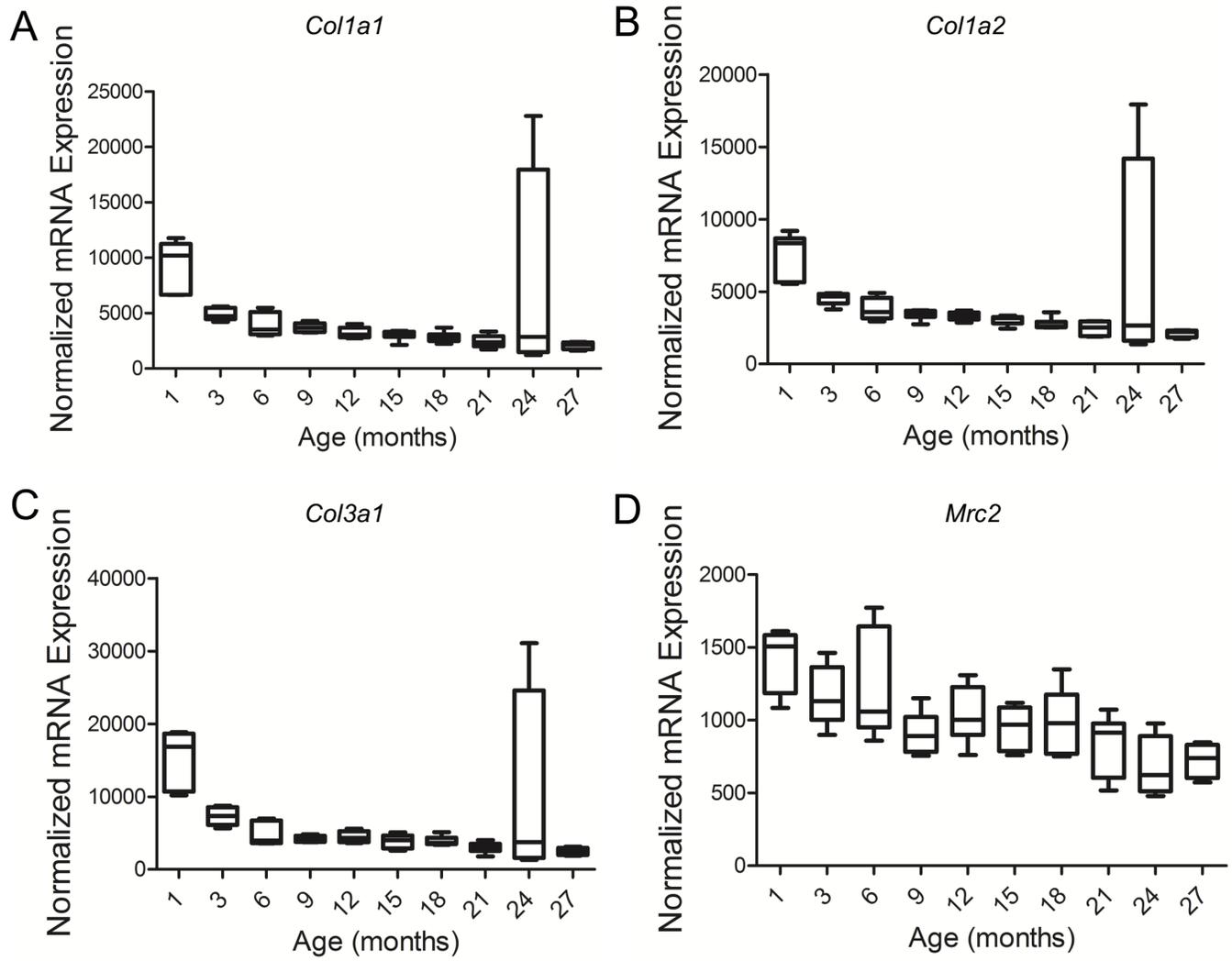
Supplemental Figure 1. (A-D) Collagen transcription and *Mrc2* transcription in whole lung decrease during the lifespan. Normalized transcriptomic data from lung over the mouse lifespan at indicated timepoints for *Mrc2* (A), *Col1a1* (B), *Col1a2* (C), *Col3a1* (D), N=3-6; data are re-analyzed from: GSE132040.

Supplemental Figure 2. *Mrc1* (Mannose receptor) transcript or protein levels are not decreased during the lifespan. (A) Representative Western blot for *Mrc1* in young versus mature whole mouse lung specimens, N=4 male mice in each group. (B) QPCR of *Mrc1* from young vs. mature mouse whole lung N=5 male mice in each group. (C) Normalized transcriptomic data from lung over the mouse lifespan for *Mrc1*.

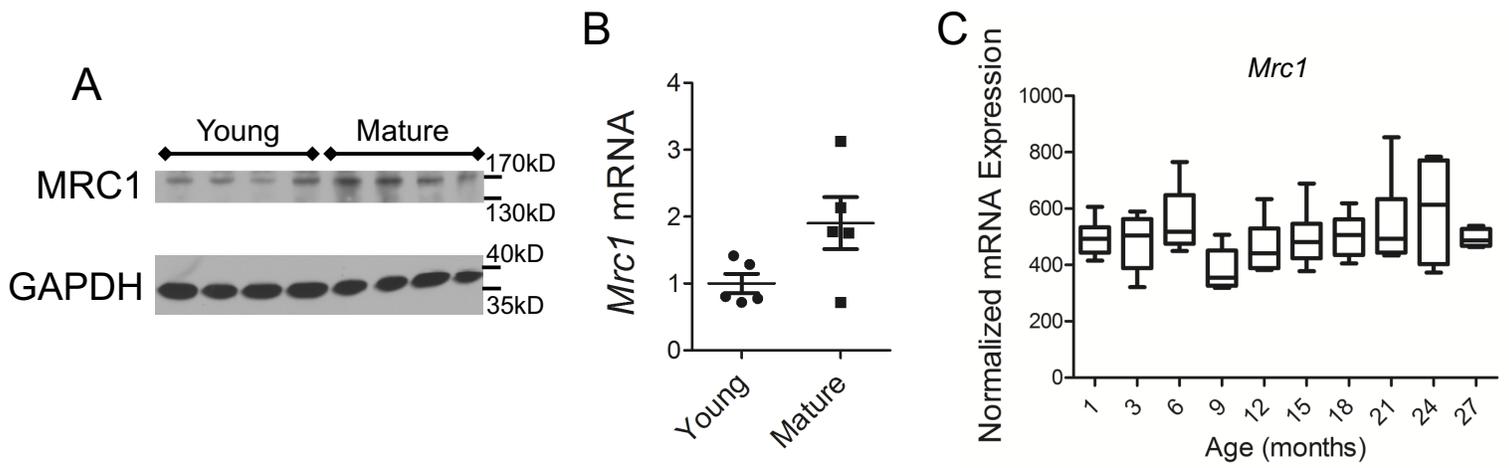
Supplemental Figure 3. Flow gating strategy. Gating strategy as pictured demonstrates gating out debris, gating for singlets, gating for live cells, gating for CD45 and then F4/80 or PDGFRA.

Supplemental Figure 4. Validation of MRC2 antibody for flow cytometric applications. (A) HEK293T cells were transfected with pCI-*Mrc2* and then stained with the sheep anti-MRC2 antibody; an untransfected control is shown as is a control without primary antibody. (B) Lysates were made from cells sorted from gates in panel (A) and then Western blot analysis was performed. (C) Flow staining of MRC2 on CD45-PDGFRA-cells is shown, N=2-4.

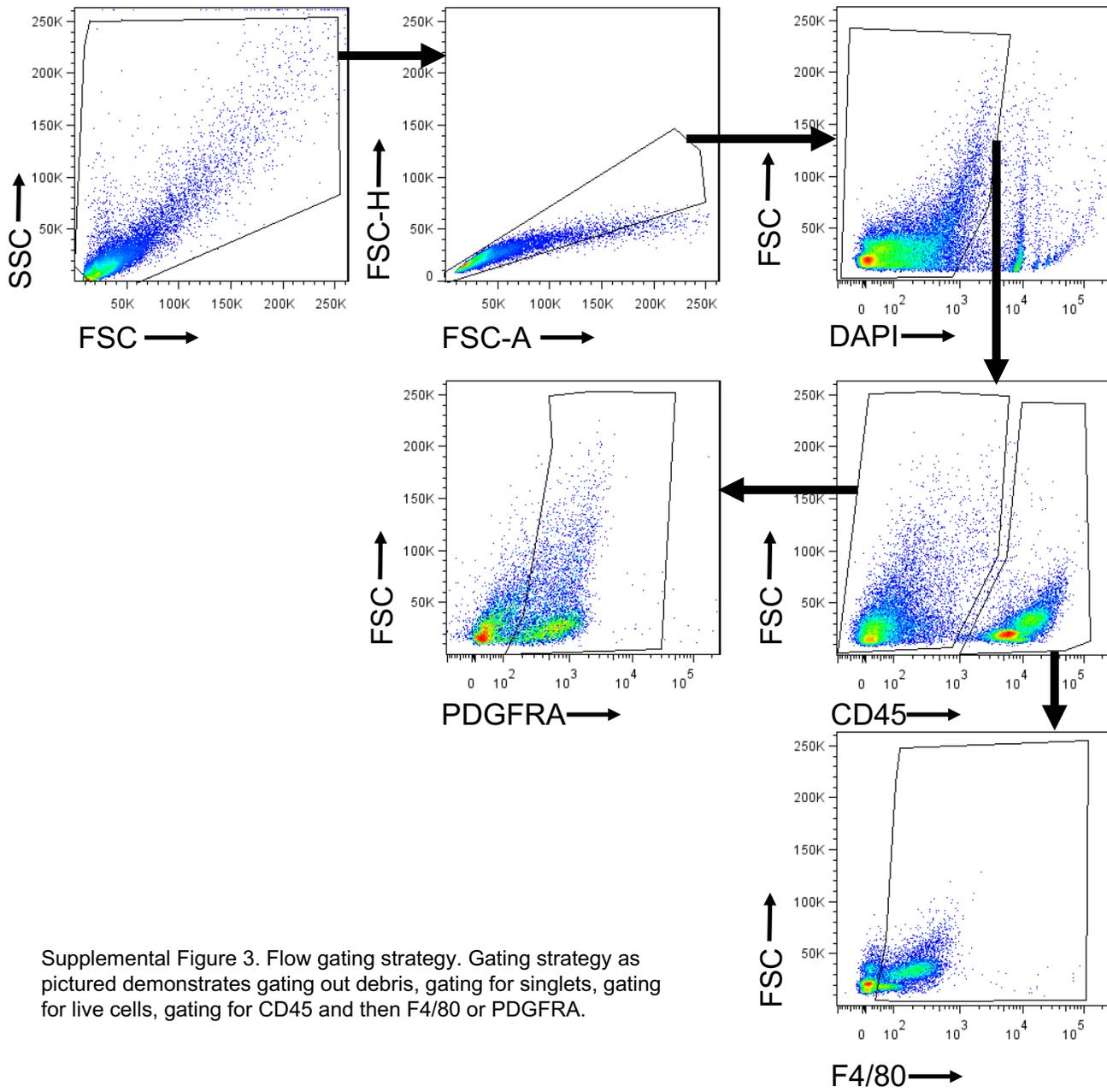
Supplemental Figure 5. Validation of *MZF1* reagents. (A) QPCR for *MZF1* after shRNA knockdown of *MZF1* in U937 cells vs. Scramble or sh-*MRC2* controls, N=5. (B) Flow cytometry for BFP in lipofectamine treated versus Empty Vector (EV, BFP only) or *MZF1*-BFP transfected cells. (C) Q-RT-PCR for *MZF1* in lipofectamine treated versus Empty vector (BFP only) or *MZF1*-BFP transfected cells. (D) Western blot of cell lysates from same conditions as (C) for *MZF1* (100kD and 129kD bands are shown, endogenous *MZF1* in U2OS cells appears at 100kD, the construct used in this figure has a 29kD BFP fused to *MZF1* giving a predicted molecular weight of 129kD); GAPDH is a loading control. Statistics: (A) ANOVA.



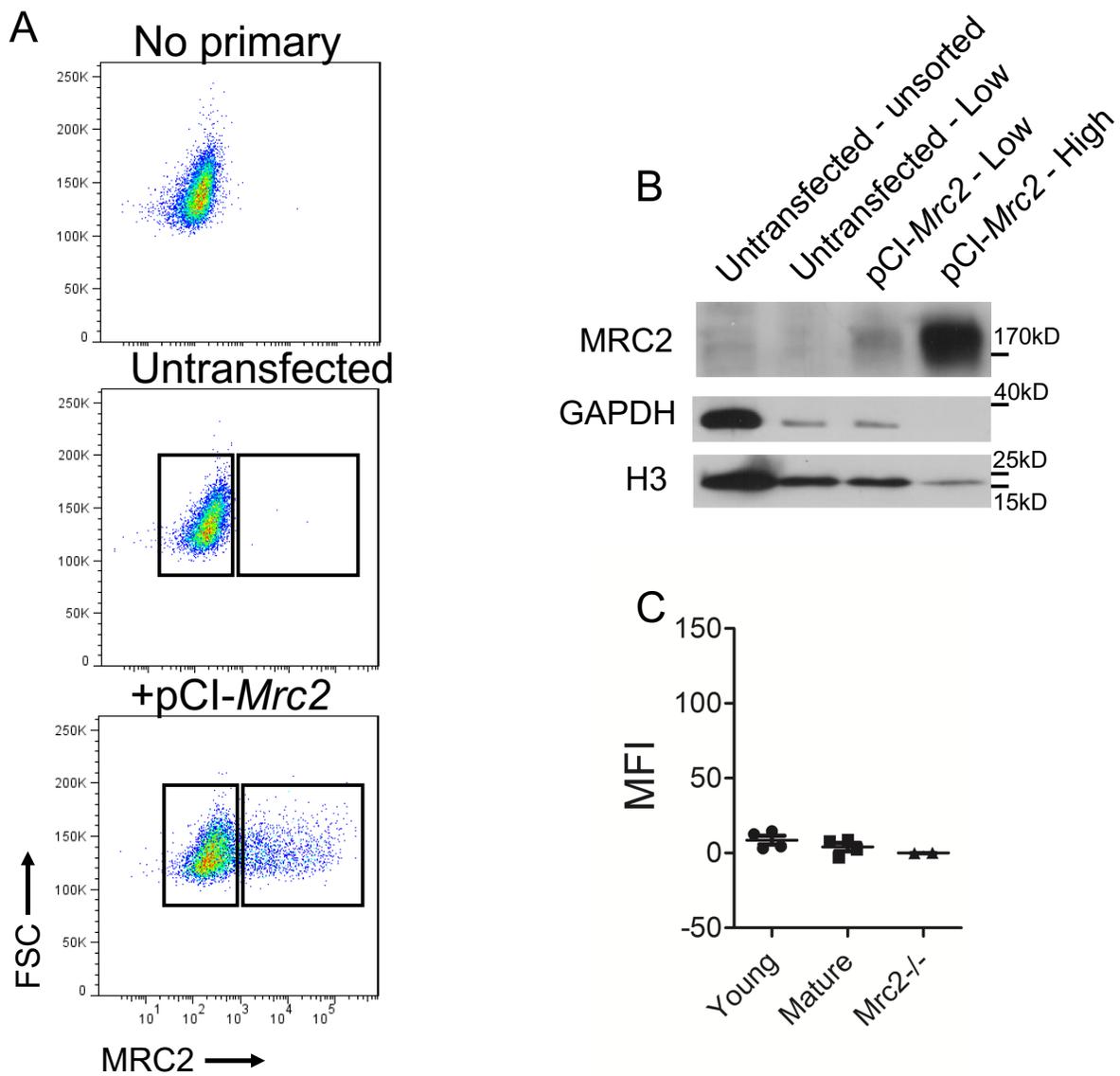
Supplemental Figure 1. (A-D) Collagen transcription and *Mrc2* transcription in whole lung decrease during the lifespan. Normalized transcriptomic data from lung over the mouse lifespan at indicated timepoints for *Mrc2* (A), *Col1a1* (B), *Col1a2* (C), *Col3a1* (D), N=3-6; data are re-analyzed from: GSE132040.



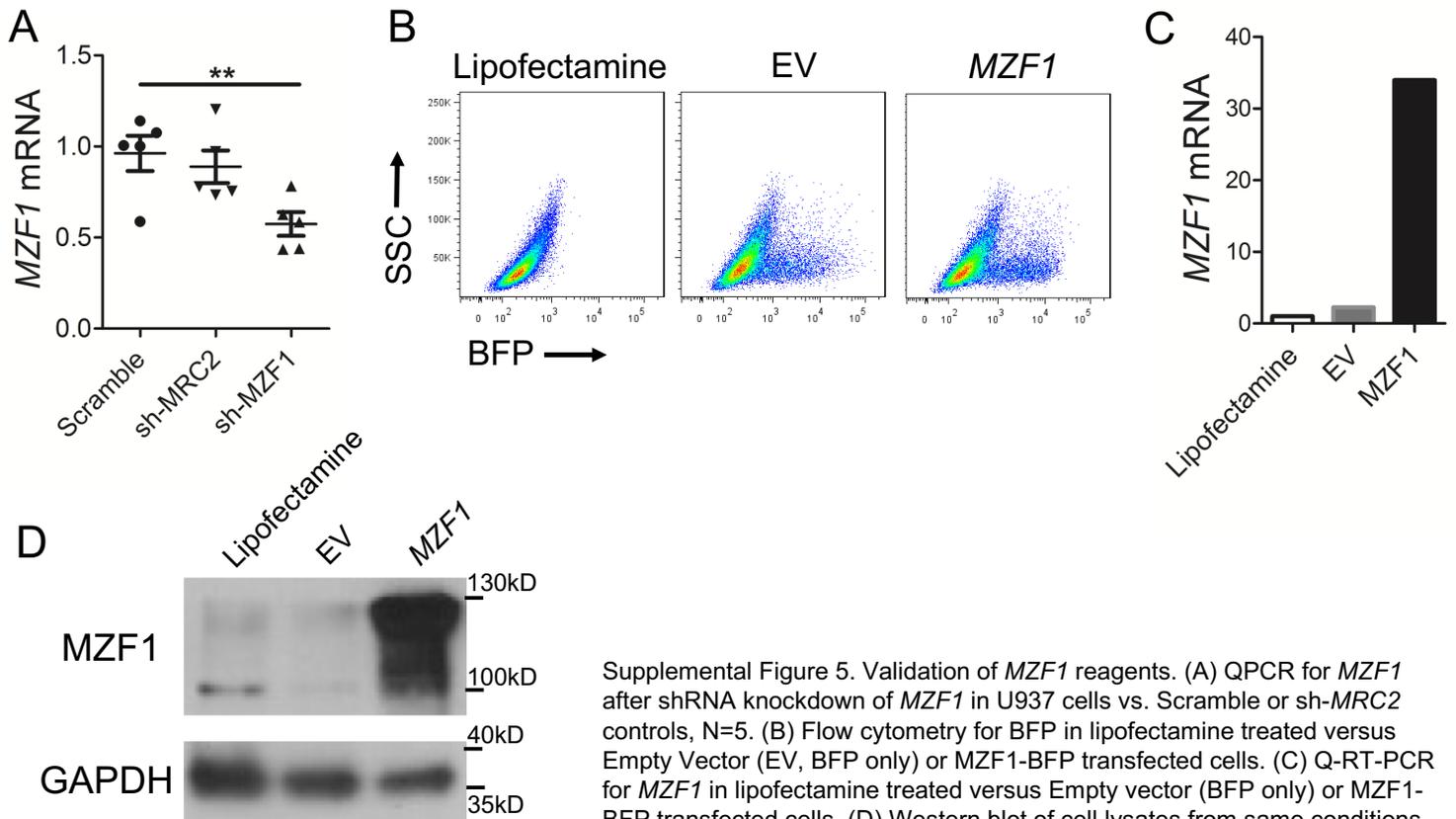
Supplemental Figure 2. *Mrc1* (Mannose receptor) transcript or protein levels are not decreased during the lifespan. (A) Representative Western blot for *Mrc1* in young versus mature whole mouse lung specimens, N=4 male mice in each group. (B) QPCR of *Mrc1* from young vs. mature mouse whole lung N=5 male mice in each group. (C) Normalized transcriptomic data from lung over the mouse lifespan for *Mrc1*.



Supplemental Figure 3. Flow gating strategy. Gating strategy as pictured demonstrates gating out debris, gating for singlets, gating for live cells, gating for CD45 and then F4/80 or PDGFRA.



Supplemental Figure 4. Validation of MRC2 antibody for flow cytometric applications. (A) HEK293T cells were transfected with pCI-*Mrc2* and then stained with the sheep anti-MRC2 antibody; an untransfected control is shown as is a control without primary antibody. (B) Lysates were made from cells sorted from gates in panel (A) and then Western blot analysis was performed. (C) Flow staining of MRC2 on CD45-PDGFR α ⁻ cells is shown, N=2-4.



Supplemental Figure 5. Validation of *MZF1* reagents. (A) QPCR for *MZF1* after shRNA knockdown of *MZF1* in U937 cells vs. Scramble or sh-*MRC2* controls, N=5. (B) Flow cytometry for BFP in lipofectamine treated versus Empty Vector (EV, BFP only) or *MZF1*-BFP transfected cells. (C) Q-RT-PCR for *MZF1* in lipofectamine treated versus Empty vector (BFP only) or *MZF1*-BFP transfected cells. (D) Western blot of cell lysates from same conditions as (C) for *MZF1* (100kD and 129kD bands are shown, endogenous *MZF1* in U2OS cells appears at 100kD, the construct used in this figure has a 29kD BFP fused to *MZF1* giving a predicted molecular weight of 129kD); GAPDH is a loading control. Statistics: (A) ANOVA.